

RAW SEQUENCE LISTING

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Application Serial Number: 10/569,792
Source: TFW
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RAW SEQUENCE LISTING

DATE: 02/15/2007

PATENT APPLICATION: US/10/569,792

TIME: 16:55:30

Input Set : A:\2007-02-14 3535-0143PUS1.ST25.txt

Output Set: N:\CRF4\02132007\J569792.raw

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3 <110> APPLICANT: ORF ehf.
4     MANTYLA, Einar
5     ORVAR, Bjorn
7 <120> TITLE OF INVENTION: A NON-DENATURING PROCESS TO PURIFY RECOMBINANT PROTEINS FROM
8     PLANTS
10 <130> FILE REFERENCE: 3535-0143PUS1
12 <140> CURRENT APPLICATION NUMBER: US 10/569,792
13 <141> CURRENT FILING DATE: 2006-02-27
15 <150> PRIOR APPLICATION NUMBER: PCT/IS04/00010
16 <151> PRIOR FILING DATE: 2004-08-27
18 <150> PRIOR APPLICATION NUMBER: 6929
19 <151> PRIOR FILING DATE: 2003-08-27
21 <160> NUMBER OF SEQ ID NOS: 3
23 <170> SOFTWARE: PatentIn version 3.3
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 564
27 <212> TYPE: DNA
28 <213> ORGANISM: Thermotoga maritima
30 <400> SEQUENCE: 1
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33 accaccgagg agatcgagac caaggccgtg gccgtgggga gcctcgacaa gaacgccacc      120
35 gccaaaggtgc gcgtgctctg ggacgagaac tacctctacg tgctcgccat cgtgaaggac      180
37 ccagtgtctca acaaggacaa cagcaacccc tgggagcaag acagcgtgga gatcttcatc      240
39 gacgagaaca accacaagac cggctactac gaggacgacg acgccaatt ccgcgtgaac      300
41 tacatgaacg agcaaacctt cgggaccggc gggagcccag cccgcttcaa gaccgccgtg      360
43 aagctcatcg aggggggcta catcgtggag gccgccatca agtgaagac catcaagcca      420
45 accccaaaca ccgtgatcgg cttcaacatc caagtgaacg acgccaacga gaaggggcaa      480
47 cgcgtgggga tcatcagctg gagcgaccca accaacaaca gctggcgcga cccaagcaag      540
49 ttcgggaacc tccgcctcat caag
52 <210> SEQ ID NO: 2
53 <211> LENGTH: 705
54 <212> TYPE: DNA
55 <213> ORGANISM: Bos taurus
57 <400> SEQUENCE: 2
58 atcgctggcg ggagcgattc cagggagggc gcatggccat gggctcgtggc actctacttc      60
60 gatgatcaac aagtctgcgg ggcattccctg gtgagcaggg attggctcgt gtccgcagca      120
62 cattgcgtgt acggcaggaa catggagcca tccaagtgga aggcagtgtc cggcctgcat      180
64 atggcatcca acctcacctc cccacaaata gagaccaggt tgatcgatca aatcgtcata      240
66 aaccacatt acaacaagcg gaggaagaac aacgacatcg caatgatgca tctcgagatg      300
68 aaggtgaact acaccgatta catacaacca atctgcttgc cagaggagaa ccaagtgttc      360
70 ccaccagggg gatctgtctc catcgcaggc tggggcgcac tcatatacca aggtccacc      420
72 gcagatgtac tgcaagaggc agacgtgcc ctcctctcca acgagaagtg ccaacaacaa      480
74 atgccagagt acaacatcac cgagaacatg gtgtgcgcag gctacgaggc aggcggggta      540

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76 gattcctgcc aaggcgattc cggcggggcca ctcatgtgcc aagagaacaa caggtggctc      600
78 ctggcaggcg tgacctcctt cggtaccacaa tgcgcactcc caaacgggcc aggggtgtac      660
80 gcacgggtgc caaggttcac cgagtggata caaagcttcc tccat                          705
83 <210> SEQ ID NO: 3
84 <211> LENGTH: 564
85 <212> TYPE: PRT
86 <213> ORGANISM: Aborophila torqueola
88 <400> SEQUENCE: 3
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91 1          5          10          15
94 Ala Cys Gly Gly Cys Ala Cys Cys Cys Cys Ala Gly Thr Gly Ala Thr
95          20          25          30
98 Cys Gly Ala Cys Gly Gly Gly Gly Ala Gly Ala Thr Cys Gly Ala Cys
99          35          40          45
102 Gly Ala Gly Ala Thr Cys Thr Gly Gly Ala Ala Cys Ala Cys Cys Ala
103          50          55          60
106 Cys Cys Gly Ala Gly Gly Ala Gly Ala Thr Cys Gly Ala Gly Ala Cys
107 65          70          75          80
110 Cys Ala Ala Gly Gly Cys Cys Gly Thr Gly Gly Cys Cys Gly Thr Gly
111          85          90          95
114 Gly Gly Gly Ala Gly Cys Cys Thr Cys Gly Ala Cys Ala Ala Gly Ala
115          100         105         110
118 Ala Cys Gly Cys Cys Ala Cys Cys Gly Cys Cys Ala Ala Gly Gly Thr
119          115         120         125
122 Gly Cys Gly Cys Gly Thr Gly Cys Thr Cys Thr Gly Gly Gly Ala Cys
123          130         135         140
126 Gly Ala Gly Ala Ala Cys Thr Ala Cys Cys Thr Cys Thr Ala Cys Gly
127 145          150         155         160
130 Thr Gly Cys Thr Cys Gly Cys Cys Ala Thr Cys Gly Thr Gly Ala Ala
131          165         170         175
134 Gly Gly Ala Cys Cys Cys Ala Gly Thr Gly Cys Thr Cys Ala Ala Cys
135          180         185         190
138 Ala Ala Gly Gly Ala Cys Ala Ala Cys Ala Gly Cys Ala Ala Cys Cys
139          195         200         205
142 Cys Cys Thr Gly Gly Gly Ala Gly Cys Ala Ala Gly Ala Cys Ala Gly
143          210         215         220
146 Cys Gly Thr Gly Gly Ala Gly Ala Thr Cys Thr Thr Cys Ala Thr Cys
147 225          230         235         240
150 Gly Ala Cys Gly Ala Gly Ala Ala Cys Ala Ala Cys Cys Ala Cys Ala
151          245         250         255
154 Ala Gly Ala Cys Cys Gly Gly Cys Thr Ala Cys Thr Ala Cys Gly Ala
155          260         265         270
158 Gly Gly Ala Cys Gly Ala Cys Gly Ala Cys Gly Cys Cys Cys Ala Ala
159          275         280         285
162 Thr Thr Cys Cys Gly Cys Gly Thr Gly Ala Ala Cys Thr Ala Cys Ala
163          290         295         300
166 Thr Gly Ala Ala Cys Gly Ala Gly Cys Ala Ala Cys Cys Thr Thr
167 305          310         315         320
170 Cys Gly Gly Gly Ala Cys Cys Gly Gly Cys Gly Gly Gly Ala Gly Cys

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171          325          330          335
174 Cys Cys Ala Gly Cys Cys Cys Gly Cys Thr Thr Cys Ala Ala Gly Ala
175          340          345          350
178 Cys Cys Gly Cys Cys Gly Thr Gly Ala Ala Gly Cys Thr Cys Ala Thr
179          355          360          365
182 Cys Gly Ala Gly Gly Gly Gly Gly Gly Cys Thr Ala Cys Ala Thr Cys
183          370          375          380
186 Gly Thr Gly Gly Ala Gly Gly Cys Cys Gly Cys Cys Ala Thr Cys Ala
187 385          390          395          400
190 Ala Gly Thr Gly Gly Ala Ala Gly Ala Cys Cys Ala Thr Cys Ala Ala
191          405          410          415
194 Gly Cys Cys Ala Ala Cys Cys Cys Cys Ala Ala Ala Cys Ala Cys Cys
195          420          425          430
198 Gly Thr Gly Ala Thr Cys Gly Gly Cys Thr Thr Cys Ala Ala Cys Ala
199          435          440          445
202 Thr Cys Cys Ala Ala Gly Thr Gly Ala Ala Cys Gly Ala Cys Gly Cys
203          450          455          460
206 Cys Ala Ala Cys Gly Ala Gly Ala Ala Gly Gly Gly Gly Cys Ala Ala
207 465          470          475          480
210 Cys Gly Cys Gly Thr Gly Gly Gly Gly Ala Thr Cys Ala Thr Cys Ala
211          485          490          495
214 Gly Cys Thr Gly Gly Ala Gly Cys Gly Ala Cys Cys Cys Ala Ala Cys
215          500          505          510
218 Cys Ala Ala Cys Ala Ala Cys Ala Gly Cys Thr Gly Gly Cys Gly Cys
219          515          520          525
222 Gly Ala Cys Cys Cys Ala Ala Gly Cys Ala Ala Gly Thr Thr Cys Gly
223          530          535          540
226 Gly Gly Ala Ala Cys Cys Thr Cys Cys Gly Cys Cys Thr Cys Ala Thr
227 545          550          555          560
230 Cys Ala Ala Gly

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VERIFICATION SUMMARY

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